IN THE CLAIMS

Please amend claims 1, 2, 5-7, 22, and 23 as follows.

- 1. (Amended) An isolated <u>corynebacterial</u> polynucleotide comprising a polynucleotide sequence selected from the group consisting of:
- a) a polynucleotide that is at least 70% identical to a polynucleotide encoding a polypeptide containing the amino acid sequence of 8EQ ID NO: 2, the polypeptide having phosphoglycerate mutase activity.
- b) a polynucleotide encoding a polypeptide containing an amino acid sequence which is at least 70% identical to the amino acid sequence of SEQ ID NO: 2, the polypeptide having phosphoglycerate mutase activity,
 - c) a polynucleotide that is complementary to the polynucleotides of a), or b), and
- d) a polynucleotide containing at least 15 consecutive bases of the polynucleotide sequence of a), b) or c), the polynucleotide encoding a polypeptide having phosphoglycerate mutase activity.
- 2. (Amended) The polynucleotide according to Claim 1 which is a DNA that replicates in corynebacterial host cells.
- 5. (Amended) An isolated corynebacterial polynucleotide comprising a polynucleotide sequence selected from the group consisting of:
- a) a polynucleotide that is at least 70% identical to a polynucleotide encoding a polypeptide containing the amino acid sequence of SEQ ID NO : 2, the polypeptide having phosphoglycerate mutase activity,
- b) a polynucleotide encoding a polypeptide containing an amino acid sequence which is at least 70% identical to the amino acid sequence of SEQ ID NO: 2, the polypeptide having phosphoglycerate mutase activity,
 - c) a polynucleotide that is complementary to the polynucleotides of a), or b), and
- d) a polynucleotide containing at least 15 consecutive bases of the polynucleotide sequence of a), b) or c), the polynucleotide encoding a polypeptide having phosphoglycerate mutase activity;

wherein the polynucleotide comprises the nucleic acid sequence as shown in SEQ ID NO: 1 and replicates in corynebacterial host cells.





- 6. (Amended) The polynucleotide that is DNA according to Claim 2 comprising:
- (i) the nucleotide sequence shown in SEQ ID NO: 1, or
- (ii) at least one sequence that is a degenerate variant of sequence (i) within the degeneracy of the genetic code], or
- (iii) the nucleotide sequence shown in SEQ ID NO: 1 in which a sense mutation has been introduced, wherein the mutated nucleotide sequence encodes for a polypeptide having phosphoglycerate mutase activity.
- 7. (Amended) An isolated corynebacterial polynucleotide comprising a polynucleotide sequence selected from the group consisting of:
- a) a polynucleotide that is at least 70% identical to a polynucleotide encoding a polypeptide containing the amino acid sequence of SEQ ID NO : 2, the polypeptide having phosphoglycerate mutase activity,
- b) a polynucleotide encoding a polypeptide containing an amino acid sequence which is at least 70% identical to the amino acid sequence of SEQ ID NO: 2, the polypeptide having phosphoglycerate mutase activity,
 - c) a polynucleotide that is complementary to the polynucleotides of a), or b), and
- d) a polynucleotide containing at least 15 consecutive bases of the polynucleotide sequence of a), b) or c), the polynucleotide encoding a polypeptide having phosphoglycerate mutase activity,

wherein the polynucleotide replicates in corynebacterial host cells and encodes a polypeptide comprising the amino acid sequence shown in SEQ ID NO: 2.

- 22. (Amended) A member of the Coryneform group of bacteria transformed by the introduction of the polynucleotide according to one of Claims 1 or 6.
- 23. (Amended) Bacteria transformed according to claim 22, wherein the bacteria are of the genus Corynebacterium.

(5 508s)

Please add following new claims 24-26.

- 24. An isolated polynucleotide isolated from the species Corynebacterium glutamicum that hybridizes to the complement of SEQ ID NO: 1, wherein the isolated polynucleotide encodes a polypeptide having the enzymatic activity of phosphoglycerate mutase.
- 25. An isolated polynucleotide comprising at least 30 consecutive nucleotides of SEQ ID NO: 1 having the function of a primer in a polymerase chain reaction to produce a polynucleotide encoding a protein comprising the amino acid sequence of SEQ ID NO: 2.
- 26. An isolated polynucleotide comprising at least 30 consecutive nucleotides of the complement to SEQ ID NO: 1 having the function of a probe in a hybridization reaction to detect or to isolate a polynucleotide encoding a protein comprising the amino acid sequence of SEQ ID NO: 2.

IN THE ABSTRACT OF THE DISCLOSURE

Please delete the present Abstract of the Disclosure and replace it with the following new Abstract of the Disclosure.

The invention provides nucleotides sequences encoding the gpm gene, which itself encodes phosphoglycerate mutase, and fermentation processes for the preparation of amino acids, especially L-lysine, using corynebacteria wherein the gpm gene is amplified.

